

PCT - Sequence listing for submission.txt.TXT  
SEQUENCE LISTING

&lt;110&gt; PERFORMANCE GENOMICS, INC.

<120> Insulin-Like Growth Factor-1 Receptor (IGF-1R) Polymorphic  
Alleles and Use of the Same to Identify DNA Markers for  
Reproductive Longevity

&lt;130&gt; 31649-2137

&lt;150&gt; US 10/662,613

&lt;151&gt; 2003-09-15

&lt;160&gt; 22

&lt;170&gt; PatentIn version 3.2

&lt;210&gt; 1

&lt;211&gt; 4500

&lt;212&gt; DNA

&lt;213&gt; Mus musculus

&lt;400&gt; 1

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PCT - Sequence listing for submission.txt.TXT

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Cys Gly Pro Gly Ile Asp Ile Arg Asn Asp Tyr Gln Gln Leu Lys Arg  
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Leu Glu Asn Cys Thr Val Ile Glu Gly Phe Leu His Ile Leu Leu Ile  
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Ser Lys Ala Glu Asp Tyr Arg Ser Tyr Arg Phe Pro Lys Leu Thr Val  
65 70 75 80

Ile Thr Glu Tyr Leu Leu Leu Phe Arg Val Ala Gly Leu Glu Ser Leu  
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Gly Asp Leu Phe Pro Asn Leu Thr Val Ile Arg Gly Trp Lys Leu Phe  
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Tyr Asn Tyr Ala Leu Val Ile Phe Glu Met Thr Asn Leu Lys Asp Ile  
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Gly Leu Tyr Asn Leu Arg Asn Ile Thr Arg Gly Ala Ile Arg Ile Glu  
130 135 140

Lys Asn Ala Asp Leu Cys Tyr Leu Ser Thr Ile Asp Trp Ser Leu Ile  
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Leu Asp Ala Val Ser Asn Asn Tyr Ile Val Gly Asn Lys Pro Pro Lys  
165 170 175

Glu Cys Gly Asp Leu Cys Pro Gly Thr Leu Glu Glu Lys Pro Met Cys  
180 185 190

Glu Lys Thr Thr Ile Asn Asn Glu Tyr Asn Tyr Arg Cys Trp Thr Thr  
195 200 205

Asn Arg Cys Gln Lys Met Cys Pro Ser Val Cys Gly Lys Arg Ala Cys  
210 215 220

Thr Glu Asn Asn Glu Cys Cys His Pro Glu Cys Leu Gly Ser Cys His  
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Thr Pro Asp Asp Asn Thr Thr Cys Val Ala Cys Arg His Tyr Tyr Tyr  
245 250 255

Lys Gly Val Cys Val Pro Ala Cys Pro Pro Gly Thr Tyr Arg Phe Glu  
260 265 270

Gly Trp Arg Cys Val Asp Arg Asp Phe Cys Ala Asn Ile Pro Asn Ala  
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275

280

285

Glu Ser Ser Asp Ser Asp Gly Phe Val Ile His Asp Asp Glu Cys Met  
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Gln Glu Cys Pro Ser Gly Phe Ile Arg Asn Ser Thr Gln Ser Met Tyr  
305 310 315 320

Cys Ile Pro Cys Glu Gly Pro Cys Pro Lys Val Cys Gly Asp Glu Glu  
325 330 335

Lys Lys Thr Lys Thr Ile Asp Ser Val Thr Ser Ala Gln Met Leu Gln  
340 345 350

Gly Cys Thr Ile Leu Lys Gly Asn Leu Leu Ile Asn Ile Arg Arg Gly  
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Asn Asn Ile Ala Ser Glu Leu Glu Asn Phe Met Gly Leu Ile Glu Val  
370 375 380

Val Thr Gly Tyr Val Lys Ile Arg His Ser His Ala Leu Val Ser Leu  
385 390 395 400

Ser Phe Leu Lys Asn Leu Arg Leu Ile Leu Gly Glu Glu Gln Leu Glu  
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Trp Asp Trp Asn His Arg Asn Leu Thr Val Arg Ser Gly Lys Met Tyr  
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Phe Ala Phe Asn Pro Lys Leu Cys Val Ser Glu Ile Tyr Arg Met Glu  
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Glu Val Thr Gly Thr Lys Gly Arg Gln Ser Lys Gly Asp Ile Asn Thr  
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Arg Asn Asn Gly Glu Arg Ala Ser Cys Glu Ser Asp Val Leu Arg Phe  
485 490 495

Thr Ser Thr Thr Thr Trp Lys Asn Arg Ile Ile Ile Thr Trp His Arg  
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Tyr Arg Pro Pro Asp Tyr Arg Asp Leu Ile Ser Phe Thr Val Tyr Tyr  
515 520 525

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Lys Glu Ala Pro Phe Lys Asn Val Thr Glu Tyr Asp Gly Gln Asp Ala  
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Cys Gly Ser Asn Ser Trp Asn Met Val Asp Val Asp Leu Pro Pro Asn  
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Lys Glu Gly Glu Pro Gly Ile Leu Leu His Gly Leu Lys Pro Trp Thr  
565 570 575

Gln Tyr Ala Val Tyr Val Lys Ala Val Thr Leu Thr Met Val Glu Asn  
580 585 590

Asp His Ile Arg Gly Ala Lys Ser Glu Ile Leu Tyr Ile Arg Thr Asn  
595 600 605

Ala Ser Val Pro Ser Ile Pro Leu Asp Val Leu Ser Ala Ser Asn Ser  
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Ser Ser Gln Leu Ile Val Lys Trp Asn Pro Pro Thr Leu Pro Asn Gly  
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Asn Leu Ser Tyr Tyr Ile Val Arg Trp Gln Arg Gln Pro Gln Asp Gly  
645 650 655

Tyr Leu Tyr Arg His Asn Tyr Cys Ser Lys Asp Lys Ile Pro Ile Arg  
660 665 670

Lys Tyr Ala Asp Gly Thr Ile Asp Val Glu Glu Val Thr Glu Asn Pro  
675 680 685

Lys Thr Glu Val Cys Gly Gly Asp Lys Gly Pro Cys Cys Ala Cys Pro  
690 695 700

Lys Thr Glu Ala Glu Lys Gln Ala Glu Lys Glu Glu Ala Glu Tyr Arg  
705 710 715 720

Lys Val Phe Glu Asn Phe Leu His Asn Ser Ile Phe Val Pro Arg Pro  
725 730 735

Glu Arg Arg Arg Arg Asp Val Met Gln Val Ala Asn Thr Thr Met Ser  
740 745 750

Ser Arg Ser Arg Asn Thr Thr Val Ala Asp Thr Tyr Asn Ile Thr Asp  
755 760 765

Pro Glu Glu Phe Glu Thr Glu Tyr Pro Phe Phe Glu Ser Arg Val Asp  
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 Val Ser Arg Gln Glu Tyr Arg Lys Tyr Gly Gly Ala Lys Leu Asn Arg  
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 Gly Asn Gly Ser Trp Thr Asp Pro Val Phe Phe Tyr Val Pro Ala Lys  
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 Thr Thr Tyr Glu Asn Phe Met His Leu Ile Ile Ala Leu Pro Val Ala  
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1270

1275

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PCT - Sequence listing for submission.txt.TXT

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PCT - Sequence listing for submission.txt.TXT

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PCT - Sequence listing for submission.txt.TXT

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PCT - Sequence listing for submission.txt.TXT

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PCT - Sequence listing for submission.txt.TXT

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PCT - Sequence listing for submission.txt.TXT

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PCT - Sequence listing for submission.txt.TXT

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PCT - Sequence listing for submission.txt.TXT

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PCT - Sequence listing for submission.txt.TXT

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PCT - Sequence listing for submission.txt.TXT

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 c 174481

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 <211> 5293  
 <212> DNA  
 <213> Sus scrofa

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 <223> n is a, c, g, or t

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PCT - Sequence listing for submission.txt.TXT

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PCT - Sequence listing for submission.txt.TXT

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tcagtggatc ttc	5293

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 <212> DNA  
 <213> Mus musculus

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<210> 9  
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 <212> DNA  
 <213> Mus musculus

PCT - Sequence listing for submission.txt.TXT

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<211> 21		
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<400> 15		
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<210> 16		
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<212> DNA		
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<400> 18		
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<211> 20		
<212> DNA		
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<400> 19		
gaagcattgg	tcggaatgta	20
<210> 20		
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<400> 21		
aagatgaggc	cttcctt	17
<210> 22		
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<212> DNA		
<213> Sus scrofa		
<400> 22		
gatcagcagg	tcgaggactg	20